## SEQUENCE LISTING

```
<110> El Tayar, Nabil
      Campbell, Robert K
      Kelton, Christie A
      He, Chaomei
<120> Novel Glycoproteins and Methods of Use Thereof
<130> 20993-003
<140> 09/927,876
<141> 2001-08-10
<150> 60/225,035
<151> 2000-08-11
<150> 60/202,724
<151> 2000-05-08
<160> 107
<170> PatentIn Ver. 2.1
<210> 1
<211> 447
<212> DÑA
<213> Homo sapiens
<400> 1
ttgaaggcag ccagatctgt taaactctgt cctttccctc tccggaagag cagcatgaag 60
ctggcattcc tetteettgg ceccatggee etecteette tggetggeta tggetgtgte 120
ctcggtgcct ccagtgggaa cctgcgcacc tttgtgggct gtgccgtgag ggagtttact 180
tteetggeea agaageeagg etgeagggge etteggatea eeaeggatge etgetggggt 240
cgctgtgaga cctgggagaa acccattctg gaacccccct atattgaagc ccatcatcga 300
gtctgtacct acaacgagac caaacaggtg actgtcaagc tgcccaactg tgccccggga 360
gtcgacccct tctacaccta tcccgtggcc atccgctgtg actgcggagc ctgctccact 420
gccaccacgg agtgtgagac catctga
                                                                   447
<210> 2
<211> 129
<212> PRT
<213> Homo sapiens
<400> 2
Met Lys Leu Ala Phe Leu Leu Leu Gly Pro Met Ala Leu Leu Leu Leu
Ala Gly Tyr Gly Cys Leu Gly Ala Ser Ser Gly Asn Leu Arg Thr Phe
                                 25
Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys Pro Gly
         35
Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg Cys Glu
                         55
Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His His
                     70
```

```
Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys Leu Pro
Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val Ala Ile
                                                     110
Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys Glu Thr
                             120
Ile
<210> 3
<211> 381
<212> DNA
<213> Xenopus sp.
<400> 3
atgaacaaga agagggtgat gttccctgtc ctgcagcttc tggttttagc cctgtgtctc 60
agcaccgctg caggatccaa tataagtctg agaacgttca ttggatgtgc tgtgagggaa 120
ttcacattct tagcaaagaa acctggctgc agaggtctgc gtgtgactac tgatgcctgc 180
tgggggcgct gtgagacctg tgagaagcca tccctagatc ctccgtacat agaagcccac 240
cacagagtet geacttacaa tgaaactaaa etggttactg taatactgee aaactgeage 300
ccagacattg acccattett tacctaccca gttgccatta gatgtgactg tgacatgtgg 360
tccacttcta ctacagaatg t
<210> 4
<211> 127
<212> PRT
<213> Xenopus sp.
<400> 4
Met Asn Lys Lys Arg Val Lys Phe Pro Val Leu Gln Leu Leu Val Leu
                                      10
Ala Leu Cys Leu Ser Thr Ala Ala Gly Ser Asn Ile Ser Leu Arg Thr
             20
Phe Ile Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys Pro
Gly Cys Arg Gly Leu Arg Val Thr Thr Asp Ala Cys Trp Gly Arg Cys
     50
Glu Thr Cys Glu Lys Pro Ser Leu Asp Pro Pro Tyr Ile Glu Ala His
                     70
His Arg Val Cys Thr Tyr Asn Glu Thr Lys Leu Val Thr Val Ile Leu
                                     90
Leu Pro Asn Cys Ser Pro Asp Ile Asp Pro Phe Phe Thr Tyr Pro Val
                                105
Ala Ile Arg Cys Asp Cys Met Trp Ser Thr Ser Thr Thr Glu Cys
                            120
<210> 5
<211> 5
```

<212> PRT

<213> Homo sapiens

<400> 5

Trp Glu Lys Pro Ile 1 5

<210> 6

<211> 141

<212> PRT

<213> Homo sapiens

<400> 6

Met Glu Met Leu Gln Gly Leu Leu Leu Leu Leu Leu Ser Met Gly

1 5 10 15

Gly Ala Trp Ala Ser Arg Glu Pro Leu Arg Pro Trp Cys His Pro Ile 20 25 30

Asn Ala Ile Leu Ala Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr
35 40 45

Val Asn Thr Thr Ile Cys Ala Gly Tyr Cys Pro Thr Met Met Arg Val 50 60

Leu Gln Ala Val Leu Pro Pro Leu Pro Gln Val Val Cys Thr Tyr Arg
65 70 75 80

Asp Val Arg Phe Glu Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val 85 90 95

Asp Pro Val Val Ser Phe Pro Val Ala Leu Ser Cys Arg Cys Gly Pro 100 105 110

Cys Arg Arg Ser Thr Ser Asp Cys Gly Gly Pro Lys Asp His Pro Leu 115 120 125

Thr Cys Asp His Pro Gln Leu Ser Gly Leu Leu Phe Leu 130 135 140

<210> 7

<211> 129

<212> PRT

<213> Homo sapiens

<400> 7

Met Lys Thr Leu Gln Phe Phe Phe Leu Phe Cys Cys Trp Lys Ala Ile 1 5 10 15

Cys Cys Asn Ser Cys Glu Leu Thr Asn Ile Thr Ile Ala Ile Glu Lys 20 25 30

Glu Glu Cys Arg Phe Cys Ile Ser Ile Asn Thr Thr Trp Cys Ala Gly

Tyr Cys Tyr Thr Arg Asp Leu Val Tyr Lys Asp Pro Ala Arg Pro Lys
50 55 60

Ile Gln Lys Thr Cys Thr Phe Lys Glu Leu Val Tyr Glu Thr Val Arg

65 70 75 80

Val Pro Gly Cys Ala His His Ala Asp Ser Leu Tyr Thr Tyr Pro Val 85 90 95

Ala Thr Gln Cys His Cys Gly Lys Cys Asp Ser Asp Ser Thr Asp Cys 100 105 110

Thr Val Arg Gly Leu Gly Pro Ser Tyr Cys Ser Phe Gly Glu Met Lys 115 120 125

Glu

<210> 8

<211> 165

<212> PRT

<213> Homo sapiens

<400> 8

Met Glu Met Phe Gln Gly Leu Leu Leu Leu Leu Leu Leu Ser Met Gly

1 5 10 15

Gly Thr Trp Ala Ser Lys Glu Pro Leu Arg Pro Arg Cys Arg Pro Ile
20 25 30

Asn Ala Thr Leu Ala Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr . 35 40 45

Val Asn Thr Thr Ile Cys Ala Gly Tyr Cys Pro Thr Met Thr Arg Val 50 55 60

Leu Gln Gly Val Leu Pro Ala Leu Pro Gln Val Val Cys Asn Tyr Arg
65 70 75 80

Asp Val Arg Phe Glu Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val 85 90 95

Asn Pro Val Val Ser Tyr Ala Val Ala Leu Ser Cys Gln Cys Ala Leu 100 105 110

Cys Arg Arg Ser Thr Thr Asp Cys Gly Gly Pro Lys Asp His Pro Leu 115 120 125

Thr Cys Asp Asp Pro Arg Phe Gln Asp Ser Ser Ser Ser Lys Ala Pro 130 135 140

Pro Pro Ser Leu Pro Ser Pro Ser Arg Leu Pro Gly Pro Ser Asp Thr 145 150 155 160

Pro Ile Leu Pro Gln 165

<210> 9

<211> 138

<212> PRT

<213> Homo sapiens

<400> 9

Met Thr Ala Leu Phe Leu Met Ser Met Leu Phe Gly Leu Ala Cys Gly .1 5 10 15

Gln Ala Met Ser Phe Cys Ile Pro Thr Glu Tyr Thr Met His Ile.Glu
20 25 30

Arg Arg Glu Cys Ala Tyr Cys Leu Thr Ile Asn Thr Thr Ile Cys Ala 35 40 45

Gly Tyr Cys Met Thr Arg Asp Ile Asn Gly Lys Leu Phe Leu Pro Lys 50 55 60

Tyr Ala Leu Ser Gln Asp Val Cys Thr Tyr Arg Asp Phe Ile Tyr Arg 65 70 75 80

Thr Val Glu Ile Pro Gly Cys Pro Leu His Val Ala Pro Tyr Phe Ser 85 90 95

Tyr Pro Val Ala Leu Ser Cys Lys Cys Gly Lys Cys Asn Thr Asp Tyr 100 105 110

Ser Asp Cys Ile His Glu Ala Ile Lys Thr Asn Tyr Cys Thr Lys Pro 115 120 125

Gln Lys Ser Tyr Leu Val Gly Phe Ser Val

<210> 10

<211> 23

<212> PRT

<213> Homo sapiens

<400> 10

Met Lys Leu Ala Phe Leu Leu Gly Pro Met Ala Leu Leu Leu Leu 1 5 10 15

Ala Gly Tyr Gly Cys Leu Gly
20

<210> 11

<211> 20

<212> PRT

<213> Homo sapiens

<400> 11

Met Glu Met Phe Gln Gly Leu Leu Leu Leu Leu Leu Leu Ser Met Gly

1 10 15

Gly Thr Trp Ala

20

<210> 12

<211> 19

<212> PRT

<213> Homo sapiens

<400> 12.

Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His

1 10 15 His Arg Val <210> 13 <211> 166 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Fusion Protein <400> 13 Met Glu Met Phe Gln Gly Leu Leu Leu Leu Leu Leu Ser Met Gly Gly Thr Trp Ala Ser Lys Glu Pro Leu Arg Pro Arg Cys Arg Pro Ile Asn Ala Thr Leu Ala Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr Val Asn Thr Thr Ile Cys Ala Gly Tyr Cys Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His His Arg Val Cys Asn Tyr 70 Arg Asp Val Arg Phe Glu Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val Asn Pro Val Val Ser Tyr Ala Val Ala Leu Ser Cys Gln Cys Ala Leu Cys Arg Arg Ser Thr Thr Asp Cys Gly Gly Pro Lys Asp His Pro 120 Leu Thr Cys Asp Asp Pro Arg Phe Gln Asp Ser Ser Ser Ser Lys Ala 135 Pro Pro Ser Leu Pro Ser Pro Ser Arg Leu Pro Gly Pro Ser Asp 160 Thr Pro Ile Leu Pro Gln 1 165 <210> 14 <211> 143 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Fusion Protein

ř, ·

10

Met Lys Leu Ala Phe Leu Leu Gly Pro Met Ala Leu Leu Leu Leu

<400> 14

Ala Gly Tyr Gly Cys Leu Gly Ala Ser Ser Gly Asn Leu Arg Thr Phe Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys Pro Gly 35 40 Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg Cys Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His His 70 Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys Leu Pro Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val Ala Ile 100 Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys Thr Val Arg Gly Leu Gly Pro Ser Tyr Cys Ser Phe Gly Glu Met Lys Glu 135 <210> 15 <211> 21 <212> PRT <213> Homo sapiens <400> 15 Cys Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala 1 5 10 His His Arg Val Cys 20 <210> 16 <211> 19 <212> PRT <213> Homo sapiens <400> 16 Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His His Arg Val <210> 17 <211> 754 <212> DNA <213> Homo sapiens <400> 17

cggcacgagg cagcaggagg cacaggaaaa ctgcaagccg ctctgttcct gggcctcgga 60 agtgatgcct atggcgtccc ctcaaaccct ggtcctctat ctgctggtcc tggcagtcac 120 tgaagcctgg ggccaggagg cagtcatccc aggctgccac ttgcacccct tcaatgtgac 180 agtgcgaagt gaccgccaag gcacctgcca gggctcccac gtggcacagg cctgtgtggg 240

```
ccactgtgag tccagcgcct tcccttctcg gtactctgtg ctggtggcca gtggttaccg 300
acacaacatc acctccgtct ctcagtgctg caccatcagt ggcctgaaga aggtcaaagt 360
acagctgcag tgtgtgggga gccggaggga ggagctcgag atcttaacgg ccagggcctg 420
ccagtgtgac atgtgtcgcc tctctcgcta ctagcccatc ctctccctc cttcctcccc 480
tgggtcacag ggcttgacat tctggtgggg gaaacctgtg ttcaagattc aaaaactgga 540
aggageteca geeetgatgg ttaettgeta tggaattttt ttaaataagg ggagggttgt 600
tecagetttg atcetttgta agattttgtg actgtcacet gagaagaggg gagtttctgc 660
ttcttccctg cctctgcctg gcccttctaa accaatcttt catcatttta cttccctctt 720
tgcccttacc cctaaataaa gcaagcagtt cttg
<210> 18
<211> 129
<212> PRT
<213> Homo sapiens
<400> 18
Met Pro Met Ala Ser Pro Gln Thr Leu Val Leu Tyr Leu Leu Val Leu
  1
                  5
Ala Val Thr Glu Ala Trp Gly Gln Glu Ala Val Ile Pro Gly Cys His
Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys
         35
Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser
Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His
65
Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys
Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu
                                105
Ile Leu Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg
                            120
Tyr
<210> 19
<211> 596
<212> DNA
<213> Mus musculus
<400> 19
eggeaegtag gggagtette agttgetgtt ggaetgteet ttgeagatge ceatggeaec 60
acgagtettg etectttgee tgetgggeet ggeagteaet gaagggeata geecagagae 120
agccatccca ggctgccact tgcacccctt caatgtgacg gtgcgcagtg atcgcctcgg 180
cacttgccag ggctcccacg tggcacaggc ctgtgtagga cactgtgagt ctagtgcttt 240
cccttcccgg tactctgtgc tggtggccag tggctatcgg cacaacatca cctcttcctc 300
ccagtgctgc accatcagca gcctcagaaa ggtgagggtg tggctgcagt gcgtggggaa 360
ccagcgtggg gagcttgaga tctttactgc aagggcctgc cagtgtgata tgtgccgttt 420
ctcccgctac tagtccccga agctcaggct ccggtcctgc cactgacatg tcatgggtat 480
ctcaaactcg gggctctgac cctctttatc gtctgtgaag atgaggttgg ccctctcagc 540
agteteettg etacattete ettegeteet gteeteaata aageaageaa tgettg
```

```
<210> 20
<211> 128
<212> PRT
<213> Mus musculus
<400> 20
Met Pro Met Ala Pro Arg Val Leu Leu Cys Leu Leu Gly Leu Ala
Val Thr Glu Gly His Ser Pro Glu Thr Ala Ile Pro Gly Cys His Leu
             20
                                 25
His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Leu Gly Thr Cys Gln
Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser Ala
Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His Asn
Ile Thr Ser Ser Ser Gln Cys Cys Thr Ile Ser Ser Leu Arg Lys Val
Arg Val Trp Leu Gln Cys Val Gly Asn Gln Arg Gly Glu Leu Glu Ile
                                105
Phe Thr Ala Arg Ile Cys Gln Cys Asp Met Cys Arg Phe Ser Arg Tyr
<210> 21
<211> 844
<212> DNA
<213> Rattus norvegicus
<400> 21
gggggaggga ggggccgaag tggccagggt tggtatgatc cccagccatg agagacatcc 60
caggggacag tgcatagaag gatggcatac acacaagtgg ctgctcattg ccttccagag 120
tagctgaggc aaggaagcaa gcaccccaca cattcccacc caaggcagag aggatcaaca 180
gtgccaccca ggcacacctc acagtcggaa gacccagaag cctggcttgc tgggggagag 240
acacaactgc aaagacttcc cttcccaccc actccttttc agatgcccat ggcacctcga 300
gtcttgctct tctgcctgct gggtctggca gtcactgaag ggcatggcct ggaggcagcc 360
gtcccaatcc caggctgcca cttgcacccc tttaacgtga cagtgcgaag tgatcgccat 420
ggcacctgcc agggctccca tgtggcacag gcgtgtgtag gacactgtga gtctagtgct 480
```

<210> 22 <211> 129

cttg

ttcccttcte ggtactctgt gctggttgce agtggctate gacacaacat cacctctgte 540 tetcagtgct gtaccatcag cagcettaaa aaggtgaggg tgtggctgca etgcgtgggg 600 aaccagegtg gggagetega gatetteaeg getagggeet gecagtgtga tatgtgeegt 660 eteteceget actaggeece gaageteagg ectecagtee tgecaetgat aggtegtget 720 teteteagae cageeetett tggagtetga agatgggget tegeetetgt ttacetggee 780 teeteageag teteaetget gettteteet teaeceetgt eeteaataaa geaggeagtg 840

1224

```
<212> PRT
<213> Rattus norvegicus
<400> 22
Met Pro Met Ala Pro Arg Val Leu Leu Phe Cys Leu Leu Gly Leu Ala
Val Thr Glu Gly His Gly Leu Glu Ala Ala Val Pro Ile Pro Gly Cys
His Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg His Gly Thr
Cys Gln Gly Ser His Val Ala Gln Ala Cys Gly His Cys Glu Ser Ser
Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His
Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Ser Leu Lys Lys
Val Arg Val Trp Leu His Cys Val Gly Asn Gln Arg Gly Glu Leu Glu
Ile Phe Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg
                            120
                                                 125
Tyr
<210> 23
<211> 1224
<212> DNA
<213> Homo sapiens
<400> 23
agatggcgaa gaaaattcca gggaagggag aatcactgca cagagggctg acacacaggt 60
cctttccaga gacagctgct cacactcaca cccatacaca cacacacaca cacacaaagg 120
cagatacagg gaaaaggcag caccattcag gcacacctca cctgtcagac cagccagccc 180
tggctcactc acctggaatg cagtatttaa agaactcgcc atcccacctg cacacccacg 240
tagagacatc tececactgt gttteagatg cetatggegt ececteaaac eetggteete 300
tatctgctgg tcctggcagt cactgaagcc tggggccagg aggcagtcat cccaggctgc 360
cacttgcacc gtgagtacct ctgggaccgg agggctagga gcagtggagg ttctgggtgg 420
gagcaaagag ctgacagagt ggacggtggg gcaggcagca ccctaaaggg ccccacactg 480
aggcacaggc aacgggagct ggggcgaggc aaaccttggc agaggcgccg tctactgctt 540
gcctatctcc ttctagcctt caatgtgaca gtgcgaagtg accgccaagg cacctgccag 600
ggeteceacg tggcacagge etgtgtggge caetgtgagt ceagegeett ceettetegg 660
tactctgtgc tggtggccag tggttaccga cacaacatca cctccgtctc tcagtgctgc 720
accatcagtg gcctgaagaa ggtgaggagg gcccgggccc ggtggatgga cgctggggtc 780
gcgggaagac cagagagatg gagatcctag acagccctga gaaaggggac tgcagcacgg 840
acteceetet eeegeaggte aaagtacage tgeagtgtgt ggggageegg agggaggage 900
tcgagatett cacggccagg gcctgccagt gtgacatgtg tcgcctctct cgctactagc 960
```

ccatcctctc ccctccttcc tcccctgggt cacagggctt gacattctgg tgggggaaac 1020 ctgtgttcaa gattcaaaaa ctggaaggag ctccagccct gatggttact tgctatggaa 1080 ttttttaaa taaggggagg gttgttccag ctttgatcct ttgtaagatt ttgtgactgt 1140 cacctgagaa gaggggagtt tctgcttctt ccctgcctct gcctggccct tctaaaccaa 1200

tctttcatca ttttacttcc ctct

```
<210> 24
 <211> 6
 <212> PRT
 <213> Homo sapiens
 <400> 24
 Leu His Pro Phe Asn Val
   1
 <210> 25
 <211> 6
 <212> PRT
 <213> Homo sapiens
<400> 25
Leu Lys Lys Val Lys Val
  1
                   5
<210> 26
<211> 116
<212> PRT
<213> Homo sapiens
<400> 26
Met Asp Tyr Tyr Arg Lys Tyr Ala Ala Ile Phe Leu Val Thr Leu Ser
Val Phe Leu His Val Leu His Ser Ala Pro Asp Val Gln Asp Cys Pro
Glu Cys Thr Leu Gln Glu Asn Pro Phe Phe Ser Gln Pro Gly Ala Pro
Ile Leu Gln Cys Met Gly Cys Cys Phe Ser Arg Ala Tyr Pro Thr Pro
     50
Leu Arg Ser Lys Lys Thr Met Leu Val Gln Lys Asn Val Thr Ser Glu
Ser Thr Cys Cys Val Ala Lys Ser Tyr Asn Arg Val Thr Val Met Gly
Gly Phe Lys Val Glu Asn His Thr Ala Cys His Cys Ser Thr Cys Tyr
                                 105
Tyr His Lys Ser
        115
<210> 27
<211> 129
<212> PRT
<213> Homo sapiens
<400> 27
Met Lys Thr Leu Gln Phe Phe Phe Leu Phe Cys Cys Trp Lys Ala Ile
                                      10
```

Cys Cys Asn Ser Cys Glu Leu Thr Asn Ile Thr Ile Ala Ile Glu Lys

20 25 30

Glu Glu Cys Arg Phe Cys Ile Ser Ile Asn Thr Trp Cys Ala Gly
35 40 45

Tyr Cys Tyr Thr Arg Asp Leu Val Tyr Lys Asp Pro Ala Arg Pro Lys
50 60

Ile Gln Lys Thr Cys Thr Phe Lys Glu Leu Val Tyr Glu Thr Val Arg 65 70 75 80

Val Pro Gly Cys Ala His His Ala Asp Ser Leu Tyr Thr Tyr Pro Val 85 90 95

Ala Thr Gln Cys His Cys Gly Lys Cys Asp Ser Asp Ser Thr Asp Cys
100 105 110

Thr Val Arg Gly Leu Gly Pro Ser Tyr Cys Ser Phe Gly Glu Met Lys 115 120 125

Glu

<210> 28

<211> 23

<212> PRT

<213> Homo sapiens

<400> 28

Met Pro Met Ala Ser Pro Gln Thr Leu Val Leu Tyr Leu Leu Val Leu

1 5 10 15

Ala Val Thr Glu Ala Trp Gly
20

<210> 29

<211> 22

<212> PRT

<213> Mus musculus

<400> 29

Met Pro Met Ala Pro Arg Val Leu Leu Cys Leu Leu Gly Leu Ala 1 5 10 15

Val Thr Glu Gly His Ser 20

<210> 30

<211> 22

<212> PRT

<213> Rattus norvegicus

<400 > 30

Met Pro Met Ala Pro Arg Val Leu Leu Phe Cys Leu Leu Gly Leu Ala 1 5 10 15

Val Thr Glu Gly His Gly

20

<210> 31

<211> 107

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus
 Sequence

<400> 31

Cys Arg Pro Gly Cys Arg Pro Thr Asn Tyr Thr Ile Ser Val Glu Lys

1 10 15

Glu Glu Cys Pro Val Cys Ile Thr Ile Asn Thr Thr Ile Cys Ala Gly
20 25 30

Tyr Cys Tyr Thr Arg Asp Pro Val Tyr Lys Ser Pro Leu Leu Pro Leu 35 40 45

Pro Gln Arg Val Cys Thr Tyr Gly Glu Trp Ser Tyr Glu Thr Ala Arg
50 55 60

Leu Pro Gly Cys Pro Pro Gly Val Asp Pro His Phe Thr Tyr Pro Val 65 70 75 80

Ala Leu Ser Cys Tyr Cys Gly Lys Cys Asn Thr Asp Thr Thr Asp Cys 85 90 95

Thr Val Leu Ser Leu Arg Pro Asp Ser Cys Ser 100 105

<210> 32

<211> 99

<212> PRT

<213> Homo sapiens

<400> 32

Thr Phe Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys

1 5 10 15

Pro Gly Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg
20 25 30

Cys Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala 35 40 45

His His Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys
50 55 60

Leu Pro Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val .65 70 75 80

Ala Ile Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys
85 90 95

Glu Thr Ile

<210> 33

<211> 107

<212> PRT

<213> Homo sapiens

<400> 33

Leu Arg Pro Arg Cys Arg Pro Ile Asn Ala Thr Leu Ala Val Glu Lys

1 5 10 15

Glu Gly Cys Pro Val Cys Ile Thr Val Asn Thr Thr Ile Cys Ala Gly
20 25 30

Tyr Cys Pro Thr Met Thr Arg Val Leu Gln Gly Val Leu Pro Ala Leu 35 40

Pro Gln Val Val Cys Asn Tyr Arg Asp Val Arg Phe Glu Ser Ile Arg 50 55 60

Leu Pro Gly Cys Pro Arg Gly Val Asn Pro Val Val Ser Tyr Ala Val 65 70 75 80

Ala Leu Ser Cys Gln Cys Ala Leu Cys Arg Arg Ser Thr Thr Asp Cys 85 90 95

Gly Gly Pro Lys Asp His Pro Leu Thr Cys Asp 100 105

<210> 34

<211> 107

<212> PRT

<213> Anguilla anguilla

<400> 34

Leu Leu Pro Cys Glu Pro Ile Asn Glu Thr Ile Ser Val Glu Lys
1 5 10 15

Asp Gly Cys Pro Lys Cys Leu Val Phe Gln Thr Ser Ile Cys Ser Gly
20 25 30

His Cys Ile Thr Lys Asp Pro Ser Tyr Lys Ser Pro Leu Ser Thr Val

Tyr Gln Arg Val Cys Thr Tyr Arg Asp Val Arg Tyr Glu Thr Val Arg
50 60

Leu Pro Asp Cys Arg Pro Gly Val Asp Pro His Val Thr Phe Pro Val 65 70 75 80

Ala Leu Ser Cys Asp Cys Asn Leu Cys Thr Met Asp Thr Ser Asp Cys
85 90 95

Ala Ile Gln Ser Leu Arg Pro Asp Phe Cys Met

<210> 35

<211> 107

<212> PRT

<213> Fundulus heteroclitus

<400> 35

Gln Leu Pro Arg Cys Gln Leu Leu Asn Gln Thr Ile Ser Leu Glu Lys 1 5 10 15

Arg Gly Cys Ser Gly Cys His Arg Val Glu Thr Thr Ile Cys Ser Gly
20 25 30

Tyr Cys Ala Thr Lys Asp Pro Asn Tyr Lys Thr Ser Tyr Asn Lys Ala 35 40 45

Ile Gln His Val Cys Thr Tyr Gly Asp Leu Tyr Tyr Lys Thr Phe Glu
50 55 60

Phe Pro Glu Cys Val Pro Gly Val Asp Pro Val Val Thr Tyr Pro Val
65 70 75 80

Ala Leu Ser Cys Arg Cys Gly Gly Cys Ala Met Ala Thr Ser Asp Cys 85 90 95

Thr Phe Glu Ser Leu Gln Pro Asp Phe Cys Met 100 105

<210> 36

<211> 109

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus
 Sequence

<400> 36

Ala Thr Lys Lys Arg Pro Lys Cys Arg Pro Thr Asn Val Thr Ile Tyr 1 15

Val Glu Lys Glu Gly Cys Thr Ser Cys Lys Thr Val Asn Thr Thr Ile 20 25 30

Cys Ala Gly Tyr Cys Tyr Thr Lys Asp Pro Val Tyr Lys Asp Gly Arg
35 40 45

Arg Leu Leu Ile Gln Cys Val Cys Cys Tyr Pro Asp Val Thr Tyr Glu
50 60

Thr Lys Val Leu Pro Gly Cys Pro Asn Gly Val Asp Pro Thr Lys Thr 65 70 75 80

Tyr Pro Val Ala Leu Ser Cys His Cys Gly Lys Cys Asn Thr Asp Asn 85 90 95

Thr Asp Cys Thr Arg Glu Ser Leu His Pro Asp Ser Cys 100 105

<210> 37

<211> 102

<212> PRT

<213> Homo sapiens

<400> 37

Asn Leu Arg Thr Phe Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu

1 5 10 15

Ala Lys Lys Pro Gly Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys
20 25 30

Trp Gly Arg Cys Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr 35 40 45

Ile Glu Ala His His Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val
50 55 60

Thr Val Lys Leu Pro Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr
65 70 75 80

Tyr Pro Val Ala Ile Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr 85 90 95

Thr Glu Cys Glu Thr Ile 100

<210> 38

<211> 109

<212> PRT

<213> Homo sapiens

<400> 38

Lys Glu Pro Leu Arg Pro Arg Cys Arg Pro Ile Asn Ala Thr Leu Ala 1 5 10 15

Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr Val Asn Thr Thr Ile 20 25 30

Cys Ala Gly Tyr Cys Pro Thr Met Thr Arg Val Leu Gln Gly Val Leu 35 40 45

Pro Ala Leu Pro Gln Val Val Cys Asn Tyr Arg Asp Val Arg Phe Glu
50 55 60

Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val Asn Pro Val Val Ser 65 70 75 80

Tyr Ala Val Ala Leu Ser Cys Gln Cys Ala Leu Cys Arg Arg Ser Thr 85 90 95

Thr Asp Cys Gly Gly Pro Lys Asp His Pro Leu Thr Cys 100 105

<210> 39

<211> 104

<212> PRT

<213> Homo sapiens

<400> 39

Asn Ser Cys Glu Leu Thr Asn Ile Thr Ile Ala Ile Glu Lys Glu Glu

1 5 10 15

Cys Arg Phe Cys Ile Ser Ile Asn Thr Ala Trp Cys Ala Gly Tyr Cys

Tyr Thr Arg Asp Leu Val Tyr Lys Asp Pro Ala Arg Pro Lys Ile Gln
35 40 45

Lys Thr Cys Thr Phe Lys Glu Leu Val Tyr Glu Thr Val Arg Val Pro 50 55 60

Gly Cys Ala His His Ala Asp Ser Leu Tyr Thr Tyr Pro Val Ala Thr
65 70 75 80

Gln Cys His Cys Gly Lys Cys Asp Ser Asp Ser Thr Asp Cys Thr Val 85 90 95

Arg Gly Leu Gly Pro Ser Tyr Cys

<210> 40

<211> 109

<212> PRT

<213> Homo sapiens

<400> 40

Arg Glu Pro Leu Arg Pro Trp Cys His Pro Ile Asn Ala Ile Leu Ala 1 5 10 15

Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr Val Asn Thr Thr Ile
20 25 30

Cys Ala Gly Tyr Cys Pro Thr Met Met Arg Val Leu Gln Ala Val Leu 35 40 45

Pro Pro Leu Pro Gln Val Val Cys Thr Tyr Arg Asp Val Arg Phe Glu
50 60

Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val Asp Pro Val Val Ser 65 70 75 80

Phe Pro Val Ala Leu Ser Cys Arg Cys Gly Pro Cys Arg Arg Ser Thr 85 90 95

Ser Asp Cys Gly Gly Pro Lys Asp His Pro Leu Thr Cys
100 105

<210> 41

<211> 109

<212> PRT

<213> Ctenolepisma lineata

<400> 41

Gly Gly Ser Leu Leu Leu Pro Cys Glu Pro Ile Asn Glu Thr Ile Ser

Val Glu Lys Asp Gly Cys Pro Lys Cys Leu Val Phe Gln Thr Ser Ile
20 25 30

Cys Ser Gly His Cys Ile Thr Lys Asp Pro Ser Tyr Lys Ser Pro Leu
35 40 45

Ser Thr Val Tyr Gln Arg Val Cys Thr Tyr Arg Asp Val Arg Tyr Glu
50 60

Thr Val Arg Leu Pro Asp Cys Arg Pro Gly Val Asp Pro His Val Thr
65 70 75 80

Phe Pro Val Ala Leu Ser Cys Asp Cys Asn Leu Cys Thr Met Asp Thr 85 90 95

Ser Asp Cys Ala Ile Gln Ser Leu Arg Pro Asp Phe Cys 100 105

<210> 42

<211> 109

<212> PRT

<213> Ctenolepisma lineata

<400> 42

Gln Ser Ser Phe Leu Pro Pro Cys Glu Pro Val Asn Glu Thr Val Ala 1 5 10 15

Val Glu Lys Glu Gly Cys Pro Lys Cys Leu Val Phe Gln Thr Thr Ile
20 25 30

Cys Ser Gly His Cys Leu Thr Lys Glu Pro Val Tyr Lys Ser Pro Phe 35 40 45

Ser Thr Val Tyr Gln His Val Cys Thr Tyr Arg Asp Val Arg Tyr Glu 50 60

Thr Val Arg Leu Pro Asp Cys Pro Pro Gly Val Asp Pro His Ile Thr 65 70 . 75 80

Tyr Pro Val Ala Leu Ser Cys Asp Cys Ser Leu Cys Thr Met Asp Thr 85 90 95

Ser Asp Cys Thr Ile Glu Ser Leu Gln Pro Asp Phe Cys. 100 105

<210> 43

<211> 109

<212> PRT

<213> Fundulus heteroclitus

<400> 43

Ala Ala Phe Gln Leu Pro Arg Cys Gln Leu Leu Asn Gln Thr Ile Ser 1 5 10 15

Leu Glu Lys Arg Gly Cys Ser Gly Cys His Arg Val Glu Thr Thr Ile
20 25 30

Cys Ser Gly Tyr Cys Ala Thr Lys Asp Pro Asn Tyr Lys Thr Ser Tyr 35 40 45

Asn Lys Ala Ile Gln His Val Cys Thr Tyr Gly Asp Leu Tyr Tyr Lys
50 60

Thr Phe Glu Phe Pro Glu Cys Val Pro Gly Val Asp Pro Val Val Thr
65 70 75 80

Tyr Pro Val Ala Leu Ser Cys Arg Cys Gly Gly Cys Ala Met Ala Thr 85 90 95

Ser Asp Cys Thr Phe Glu Ser Leu Gln Pro Asp Phe Cys 100 105

<210> 44

<211> 105

<212> PRT

<213> Rana catesbeiana

<400> 44

Arg His Val Cys His Leu Ala Asn Ala Thr Ile Ser Ala Glu Lys Asp 1 5 10 15

His Cys Pro Val Cys Ile Thr Phe Thr Thr Ser Ile Cys Thr Gly Tyr
20 25 30

Cys Gln Thr Met Asp Pro Val Tyr Lys Thr Ala Leu Ser Ser Phe Lys 35 40 45

Gln Asn Ile Cys Thr Tyr Lys Glu Ile Arg Tyr Asp Thr Ile Lys Leu 50 55 60

Pro Asp Cys Leu Pro Gly Thr Asp Pro Phe Phe Thr Tyr Pro Val Ala 65 70 75 80

Leu Ser Cys Tyr Cys Asp Leu Cys Lys Met Asp Tyr Ser Asp Cys Thr 85 90 95

Val Glu Ser Ser Glu Pro Asp Val Cys . 100 105

<210> 45

<211> 111

<212> PRT

<213> Anguilla anguilla

<400> 45

Ala Gly Gln Val Leu Ser Ile Cys Ser Pro Val Asp Tyr Thr Leu Tyr

1 10 15

Val Glu Lys Pro Glu Cys Asp Phe Cys Val Ala Ile Asn Thr Thr Ile 20 25 30

Cys Met Gly Phe Cys Tyr Ser Leu Asp Pro Asn Val Val Gly Pro Ala . 35 40 45

Val Lys Arg Leu Val Val Gln Arg Gly Cys Thr Tyr Gln Ala Val Glu 50 60

Tyr Arg Thr Ala Glu Leu Pro Gly Cys Pro Pro His Val Asp Pro Arg 65 70 75 80

Phe Ser Tyr Pro Val Ala Leu His Cys Thr Cys Arg Ala Cys Asp Pro 85 90 95

Ala Arg Asp Glu Cys Thr His Arg Ala Ser Ala Asp Gly Asp Arg

100 105 110

<210> 46 <211> 23 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR Primer Sequence <400> 46 tcgatgatgg gcttcaatat agg 23 <210> 47 <211> 26 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR Probe Sequence <400> 47 cctgggagaa acccattctg gaaccc 26 <210> 48 <211> 22 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR Primer Sequence <400> 48 gcctcagatg gtctcacact cc 22 <210> 49 <211> 29 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: PCR Primer Sequence <400> .49 ctcgaggcct ccagtgggaa cctgcgcac 29 <210> 50 <211> 31 <212> DNA <213> Artificial Sequence <220>

<223>	Description of Artificial Sequence: PCR Primer Sequence	
<400> gggcc	50 cggat cetcagatgg tetcacaete e	31
<210>	51	
<211>	· · · · · · · · · · · · · · · · · · ·	
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Description of Artificial Sequence: PCR Primer Sequence	
<400>	51	
-	catga agctggcatt cctc	24
-210s		
<210><211>		
<212>	<del></del>	
	Artificial Sequence	
	-	
<220>		
<223>	Description of Artificial Sequence: PCR Primer Sequence	
	bequence	
<400>	52	
tatcga	atggt ctcacactcc gtg	23
<210>	53	
<211>		
<212>	DNA	
<213>	Artificial Sequence	
<220>		
	Description of Artificial Sequence: Synthetic	
12857	Oligonucleotide	
	•	
<400>		
ctagto	ctcga ggctgcagtt gctgactaca aagacgatga cgacaagg	48
<210>	54	
<211>		
<212>		
<213>	Artificial Sequence	
.0.0.5		
<220>	Doggwinking of Autofal A a	
<b>\</b> 443>	Description of Artificial Sequence: Synthetic Oligonucleotide	
<400>		
ccttgt	cgtc atcgtctttg tagtcagcaa ctgcagcctc gaga	44
<210>	55	
<211>		

<212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: PCR Primer Sequence	
<400> 55 tttgctagca ccatgtctgc acttctg	27
<210> 56 <211> 27 <212> DNA <213> Artificial Sequence	
<pre>&lt;220&gt; &lt;223&gt; Description of Artificial Sequence: PCR Primer</pre>	
<400> 56 tttggatcet cagatggtet caeacte	27
<210> 57 <211> 19 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: PCR Primer Sequence	
<400> 57 aggaggcagt catcccagg	19
<210> 58 <211> 18 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: PCR Primer Sequence	
<400> 58 tgccttggcg gtcacttc	18
<210> 59 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: PCR Probe Sequence	
<400> 59	24

```
<210> 60
  <211> 17
  <212> DNA
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: PCR Primer
        Sequence
  <400> 60
  aggcagccgt cccaatc
                                                                      17
  <210> 61
  <211> 22
  <212> DNA
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: PCR Primer
        Sequence
 <400> 61
 gatcacttcg cactgtcacg tt
                                                                      22
 <210> 62
 <211> 22
 <212> DNA
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: PCR Probe
       Sequence
 <400> 62
 caggctgcca cttgcacccc tt
                                                                      22
 <210> 63
 <211> 31
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: PCR Primer
       Sequence
 <400> 63
 ttttaagctt agtgatgcct atggcgtccc c
                                                                      31
<210> 64
 <211> 25
 <212> DNA
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: PCR Primer
```

·z: . `

Sequence <400> 64 ttttgaattc gtagcgagag aggcg 25 <210> 65 <211> 22 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: PCR Primer Sequence <400> 65 tttgagatct tcacggccag gg 22 <210> 66 <211> 17 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Oligonucleotide <400> 66 ctagaggaat tcgggcc 17 <210> 67 <211> 9 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Oligonucleotide <400> 67 cgaattcct 9 <210> 68 <211> 31 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR Primer Sequence <400> 68 ttttctagaa caggaggcag tcatcccagg c 31

<210> 69 <211> 28 <212> DNA

```
<213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: PCR Primer
       Sequence
 <400> 69
ttttgaattc ctagtagcga gagaggcg
                                                                     28
<210> 70
 <211> 52
 <212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 70
agttgctgac tacaaagacg atgacgacaa gcaggaggca gtcatcccag gc
                                                                   . 52
<210> 71
<211> 42
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 71
cccgtttaaa cggatcctca gtagcgagag aggcgacaca tg
                                                                     42
<210> 72
<211> 74
<212> DNA
<213> Artificial Sequence
<22.0>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 72
tttgctagcc accatgtctg cacttctgat cctagctctt gttggagctg cagttgctga 60
ctacaaagac gatq
<210> 73
<211> 42
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
cccgtttaaa cggatcctca gtagcgagag aggcgacaca tg
                                                                    42
```

```
<210> 74
 <211> 26
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: PCR Primer
       Sequence
 <400> 74
ggtaccaagg tagccttgca gaagtt
                                                                     26
<210> 75
 <211> 26
 <212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
       Sequence
<400> 75
cagctggtaa ttgaactggg agtgga
                                                                     26
<210> 76
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 76
gggccttcgg atcaccac
                                                                     18
<210> 77
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 77
cagcatgaag ctggcattcc tc
                                                                    22
<210> 78
<211> 6240
<212> DNA
<213> Homo sapiens
<400> 78
aggaatetet ggatgeetgt gttggagttt gtgggeattt acaatttetg ggeteatttt 60
```

ccctgaaatg ctaggagcaa ggtccctttg atagtgacaa atgcatggtt ggctgtgcca 120 ttgaaggcag ccagatctgt taaactctgt cctttccctc tccggaagag cagcatgaag 180 ctggcattcc tetteettgg ceccatggcc etceteette tggctggcta tggctgtgtc 240 ctcggtgcct ccagtgggaa cctgcgcacc tttgtgggct gtgccgtgag ggagtttact 300 ttcctggcca agaagccagg ctgcaggggc cttcggatca ccacggatgc ctgctggggt 360 cgctgtgaga cctgggaggt gagttgctaa gttgtgcaga tgacagtgtc ttctaggcca 420 gcagcttggg tctgattctt aagagttcac tttttaaatg atatgaggta gagctgggac 480 agtgatttga aaaacatgat gttgcccctc taacaaagca ttgataaggt taagaatttg 600 gtttacattg tgtctatgta tctgggaatc atctctggga ggtcaagatg tactgttcta 660 cccgttttac agatgacatg gagggattca agggagagtg gctgcaaagt cacgtagagc 720 gtcagtgtaa agctgggaat caatctgtgg ttcaagcttg tgacccaaac tcctccctat 780 gtttcctcat tttggataaa ttagccagtt tccaagaaag aggccctgag ctgaagggtg 840 agcgttggtc ccagtgaagg gtgagacccc ttcactgcct cttctgcagc ccttttcctc 900 ctcaagtctc tgggagccct ctggggttat cactgacgga tccattaagt tccttcatat 960 tcaattatac ctggcctttt tagagacatt taatttaaag tggagataac actctcaaac 1020 aaagttaaaa teetattggg etaagaggag etgtttgagt gatgaagagg aagagageta 1080 ttcagcaccc cagcagatca cattacgtag tgactgtggg ctcttccccc tgaggcctgc 1140 ccacttggta accaatgaag tgctgtctct gatcttgtca ctccctggcc caaaaacctt 1200 gaatgtccac acactactac agattcaata actaactttc aaggtgctca gcaatatggc 1260 gtctgcctgc tttcctggag acagcacatt ttcttactct ggccttggta agtgactttc 1320 aaaggtttta tcaaatagcc cttatggatc tcattttgtt ccttccctca tatcccttct 1380 ccttcccatc tgtcattatc atatttattc ctgatgccta tctgcagtgc cagctccctt 1440 tctgggcctt ttttgacttg caggtaagcc cttgactatg ctctactttt cgtcttactt 1500 cctccccac cacacgcgtg atttaaattt tttcaggaca gaggttcatt cttataacct 1560 tcacagcttt tgtcaagatg tcgtgtatga acaaggcatt caatacacat ttgttggttg 1620 actgggatgg acctcccct ggagctgtag atcctccagc ctaatggaag gccatttaga 1680 atcacacttg cactgtgagt ggacactgcc attgggaaaa atagccttct ctttggggac 1740 ccagagggta acctgetett gettaggtae aattaeggee etgtgaatgg aattgggtea 1800 tagtgatgaa atctccaaat tggatgaaac tactctatca aagtagtttt cttttgcctc 1860 attcaggggc ttgagcccta ctagcccaat gaaaatcggg ttttgctaag tagactttgc 1920 ctgtcaattg gcagcaaatt cacctggggc acttggcacc tcctcctgtt cagggactgg 1980 cctggcaggg cctctccctg ttcgcatcta gtgtctgggc tatttgaagc cctctctgtg 2040 tgatgaatgt ctttaattgg atcatggtca cccataggag gtcaggaact gtgctctcac 2160 tggaaagatg gaaacaccaa aaccgttaaa gaacaagatt ctccctgatg ttagccagct 2220 ttcattcatg tcttgactgt gttatgaaaa gggaggttac ctatagaaaa taaataaaag 2280 aatgagattc attttcccag caatctgaaa gtttctgcgc tataaagcac ttgattttt 2340 ggtgggggg atcttaactg aaagcatgtc tgaaaataag gatgttcatg atgacaggct 2400 ggctggattt acatttgaag gttgttgaaa atagctattc ctcataatct gggtatagag 2460 ttgccagatt tagcaaacaa acaaacagac aaacaaaata aaacaaaacc aatcccctcc 2520 ccacagaaac ccaaactgaa ataaaaccag aaaaccagga agcccaggta aattggaatt 2580 taagataaat aataaataaa tttttagcgt aagtctgtct gtctcataca gtatttggga 2640 tgacttatac taaaaaatta tgtatctgaa aatgaaattt tacggggcgt ttggtctgcc 2700 taggttccca gagtactaat ggtaagagga cttaaagcaa atacgggaag gtaggagaaa 2760 acagttcagg acaaattcag ctcttctggt ctttgtcaaa ggcaaggctg gccgggcgtg 2820 gtggctaaca cctgtaatct cagcactttg ggaggctgtg gtgggtggat aatgaggtca 2880 ggagttcgag accagcctgg ccagttttta gtaaagaggt gagttaaacc ctgtctctac 2940 taaaaataca aaaattagcc gggcatggtg gtatgcacct gtagtcccag ctacttggga 3000 ggctgaggca gaagacttgc ttgaacccag gaggtggagg ttacagtgag ccaagatcat 3060 gccactatac tccagcctgg cgacagagtg agactccatc tcaaaaaaaa aaaaaaaga 3120 aaaaagaaaa aaaaaaggta aggctgctat tttcatgaca ttcatgcaag aacatcttga 3180 gttacatatg tatatatt cttttttgcc tagaacaaag aagaaccaaa aagcaaaggt 3240 actgtcattt gaaagcttgt tattatttac attactttct tataataatt gcactaataa 3300 gaacaatgga ttggctgggc gtggtggctc acgcctgtaa tcccagcact ttgggaggcc 3360 gaggcaggca gatcacgagg tcaggaaatc gagaccatcc tggctaacat ggtgaaaccc 3420 tgtctctact aaaaatacaa aaaatgagcc aggcgtggtg gtgggtgcct gtagtcccgg 3480 gaggctgagg caggagaatg gcgtgaaccc gggaggcgga gattgcaatg agctgagatt 3540 gcgccactga actccagcct gggagacagc aagactccgt ctcaaaaaaa aaaaaaatgg 3600 attgcatttt ttgaacattt actttgttct agacattgtg cattgcgtat atcatcttac 3660 cttatctctc aaacaatggt gggaggtagc tattttgttt tacagaggag gaaacttgag 3720

```
tetteaggaa gttaagtgga tttteeaagg tetecageaa gtggeagaac agggaeteaa 3780
  gctccttagt tctgactgca gggctcgaga ttttaactcc agctaggtgc tgatattttt 3840
  tctgatctgt gtgttctgtt tatcaaaatt gtctttgaac ttaagattta taaaaggtga 3900
  aggaaggaaa tgaatctttt tgatgatcag aacagtgcac agagtattcg ggaacctgtc 3960
  ttgtaatgtt ttctttcatt gattcaatga caaatagtta ttgaaactct cccggggtct 4020
  gttttgggta cttgaggcac agtgggcaaa aatctctgtc ctaaaagagc ttactttcta 4080
  gagtgggagg aatatcacac gaatgaaagg tagactacgt cgtgtggtat tgatcagtgc 4140
  tgtggtggaa aataaagcaa gatgggggat gggaagtttc tgggcatgga gatggaatgt 4200
  tgcaatttta aataggatgg tcaggaaatg cttccctgag agggtgacat tctaacaaaa 4260
  acccaaggtt ggtgaaagag tgaatcatac gggagaagaa tgttccaggc agaaggaacg 4320
  gtaagtgcaa aggccctgag ctggggctgt tcctggtggg tcagaggagc aataaggaga 4380
  ccgccgtgag cctagtgagg aagtcagtga ggtgggaatg gttgcaggca tttcagaagg 4440
  tagagttgca gagaaggtga tgtaggtctt gaaggtgatc ataaggtctt tgatgtttgt 4500
  tctgagtgag atgggaaatc actggggctt tgggcagagg agtgacatga tctgacttag 4560
  gtttaaacag gatcactcag ggccgctgtg ttgcaaatag attgtaggga gtaaaaatgg 4620
  aagaggggag accagttaga aggtatttgc aatgactaag atgattcatt tgctgactat 4680
  gcatggagca cttgctgtgt gctatggtct ctcctgggag cttagaatat ggtcttgagt 4740
  gaaatcagct tcttgctttc aggagtttgt tttctactgg gagacgacag agcaacaagt 4800
  aaatcaacga ataacaagtt aatttctgat agtgataaat gatactaaaa aactgaaaca 4860
  agatcatatg ttctaatgaa ttctctgttt tctatctatg gggacagaaa cccattctgg 4920
  aaccccccta tattgaagcc catcatcgag totgtaccta caacgagacc aaacaggtga 4980
  ctgtcaagct gcccaactgt gccccgggag tcgacccctt ctacacctat cccgtggcca 5040
  tccgctgtga ctgcggagcc tgctccactg ccaccacgga gtgtgagacc atctgaggcc 5100
  gctagctgct ctctgcagac ccacctgtgt gagcagcaca tgcagttata cttcctggat 5160
  gcaagactgt ttaatttcga ccacacccat ggaggaggtt acctgtcgcc ccttaggtcc 5220
  ageteaggea aaaggeeeaa atgeageeta ettatgetaa aagtteaaaa caatattegt 5280
  gccttcacca aaataatttc tccagctcac atacctgcaa attaattttt ctttgccttg 5340
  agtcttggaa cataatttgt gtatcacaat cctcccccaa tttggactta taatatgcta 5400
  atgatttaaa cacatgggat gtaattagga tatggggctg gaaagtcttt aaattctcat 5460
  gttctattta acctctgatc tccaaccgga tttatgatta aagggctaga aatgaacaaa 5520
  acccatgtac tagtetteet taccccagag gaattecage tgcaagette tttagggaaa 5580
  atgetecett eccettttaa etgageaatt atetacaeaa gaaataagae tgeteagata 5640
  tacaaagaga gtagcttcaa tgaaaagatg tttggatttg gataattctt ttccctagca 5700
  aaattegeta geteeettaa gagtettaat aaagaggeta egttgggatt aaaagaaaaa 5760
  aaaacagaaa taaaatatgt aactaatagc tatctcattt agccttaaaa acttattaaa 5820
  ctaaactcat gttttagagt atgatgttct cccaaagcta tggcaaaatg gccaatcaca 5880
 agtattette eccatttate atatttteaa tttaagttgt aaettaetaa aeteagaaat 5940
 .tttatatgcg tttaggggta aaactgcatg gctggctcag aggaaaaagc ctgtgatttt 6000
 ctageteetg cetetetaaa atettacagt agetaattet gtggetggaa aaaaceteea 6060
 aaactctaat gttatgcaaa tgtctttaat tctggcattt ttggggttga atttaacctt 6120
 gttccttttt cataatgtgc caagaaaacc tatattaatg ccaataaagc atgtcctctg 6180
 tcttttggat tcatgacaac attcaagaaa gtctttttaa ttcttagtat acttggagta 6240
 <210> 79
 <211> 1224
 <212> DNA
 <213> Homo sapiens
 <400> 79
 agatggcgaa gaaaattcca gggaagggag aatcactgca cagagggctg acacacaggt 60
 cctttccaga gacagctgct cacactcaca cccatacaca cacacacaca cacacaaagg 120
 cagatacagg gaaaaggcag caccattcag gcacacctca cctgtcagac cagccagccc 180
 tggctcactc acctggaatg cagtatttaa agaactcgcc atcccacctg cacacccacg 240
 tagagacatc tececactgt gttteagatg cetatggegt ececteaaac eetggteete 300
 tatetgetgg teetggeagt caetgaagee tggggecagg aggeagteat eecaggetge 360
^\prime cacttgcacc gtgagtacct ctgggaccgg agggctagga gcagtggagg ttctgggtgg 420 ^\prime
 gagcaaagag ctgacagagt ggacggtggg gcaggcagca ccctaaaggg ccccacactg 480
 aggcacaggc aacgggagct ggggcgaggc aaaccttggc agaggcgccg tctactgctt 540
 gcctatctcc ttctagcctt caatgtgaca gtgcgaagtg accgccaagg cacctgccag 600
 ggctcccacg tggcacaggc ctgtgtgggc cactgtgagt ccagcgcctt cccttctcgg 660
```

```
tactctgtgc tggttgccag tggttaccga cacaacatca cctccgtctc tcagtgctgc 720
    accatcagtg gcctgaagaa ggtgaggagg gcccgggccc ggtggatgga cgctggggtc 780
    gcgggaagac cagagagatg gagatcctag acagccctga gaaaggggac tgcagcacgg 840
    actococtot coogcaggto aaagtacago tgcagtgtgt ggggagcogg agggaggago 900
    tcgagatctt cacggccagg gcctgccagt gtgacatgtg tcgcctctct cgctactagc 960
    ccatcctctc ccctccttcc tcccctgggt cacagggctt gacattctgg tgggggaaac 1020
    ctgtgttcaa gattcaaaaa ctggaaggag ctccagccct gatggttact tgctatggaa 1080
    tttttttaaa taaggggagg gttgttccag ctttgatcct ttgtaagatt ttgtgactgt 1140
    cacctgagaa gaggggagtt tetgettett ceetgeetet geetggeeet tetaaaccaa 1200
    tctttcatca ttttacttcc ctct
   <210> 80
   <211> 490
   <212> DNA
   <213> Artificial Sequence
   <220>
   <223> Description of Artificial Sequence: Fusion Protein
  <400> 80
. Cactttgcct ttctctccac aggtgtccac tcccagttca attaccagct gctagcgtcg 60
accatgtctg cacttctgat cctagetett gttggagetg cagttgetea teatcaccat 120
  , caccatggtg acgatgacga taagcaggag gcagtcatcc caggctgcca cttgcacccc 180
ttcaatgtga cagtgcgaag tgaccgccaa ggcacctgcc agggctccca cgtggcacag 240
 gcctgtgtgg gccactgtga gtccagcgcc ttcccttctc ggtactctgt gctggtggcc 300
   agtggttacc gacacaacat cacctccgtc tctcagtgct gcaccatcag tggcctgaag 360
aaggtcaaag tacagetgca gtgtgtgggg ageeggaggg aggagetega gatetteaeg 420
es gccagggcct gccagtgtga catgtgtcgc ctctctcgct actagtcgac ggatccagac 480
atgataagat
                                                                     490
: <210> 81
on <211> 130
  <212> PRT
   <213> Homo sapiens
   <4.00> 81
... Met Lys Leu Ala Phe Leu Phe Leu Gly Pro Met Ala Leu Leu Leu
Ala Gly Tyr Gly Cys Val Leu Gly Ala Ser Ser Gly Asn Leu Arg Thr
                                    25
   Phe Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys Pro
   Gly Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg Cys
   Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His
                        70
   His Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys Leu
  Pro Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val Ala
              100
                                                       110
  Ile Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys Glu
          115
                                                   125
```

```
Thr Ile
     130
 <210> 82
 <211> 420
 <212> DNA
 <213> Homo sapiens
 <400> 82
 cgaattcgcc cttcagcatg aagctggcat tcctcttcct tggccccatg gccctcctcc 60
 ttdtggetgg etatggetgt gteeteggtg eeteeagtgg gaacetgege acetttgtgg 120
 getgtgeegt gagggagttt aettteetgg ceaagaagee aggetgeagg ggeettegga 180
 tcaccacgga tgcctgctgg ggtcgctgtg agacctggga gaaacccatt ctggaaccc 240
 cctatattga agcccatcat cgagtctgta cctacaacga gaccaaacag gtgactgtca 300
 agctgcccaa ctgtgccccg ggagtcgacc ccttctacac ctatcccgtg gccatccgct 360
 gtgactgcgg agcctgctcc actgccacca cggagtgtga gaccatctga ggcaagggcg 420
 <210> 83
 <211> 106
 <212> PRT
 <213> Artificial Sequence
 <220> ...
 <223> Description of Artificial Sequence: Fusion Protein
<400> .83
Ala Ser Ser Gly Asn Leu Arg Thr Phe Val Gly Cys Ala Val Arg Glu
Phe Thr Phe Leu Ala Lys Lys Pro Gly Cys Arg Gly Leu Arg Ile Thr
Thr Asp Ala Cys Trp Gly Arg Cys Glu Thr Trp Glu Lys Pro Ile Leu
Glu Pro Pro Tyr Ile Glu Ala His His Arg Val Cys Thr Tyr Asn Glu
Thr Lys Gln Val Thr Val Lys Leu Pro Asn Cys Ala Pro Gly Val Asp
Pro Phe Tyr Thr Tyr Pro Val Ala Ile Arg Cys Asp Cys Gly Ala Cys
                 85
Ser Thr Ala Thr Thr Glu Cys Glu Thr Ile
            100
<210> 84
<211> 420
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Fusion Protein
<400> 84
ggactagtcc tgcaggttta aacgaattcg cccttctcga ggcctccagt gggaacctgc 60
```

```
gcacctttgt gggctgtgcc gtgagggagt ttactttcct ggccaagaag ccaggctgca 120
 ggggccttcg gatcaccacg gatgcctgct ggggtcgctg tgagacctgg gagaaaccca 180
 ttctggaacc cccctatatt gaagcccatc atcgagtctg tacctacaac gagaccaaac 240
aggtgactgt caagctgccc aactgtgccc cgggagtcga ccccttctac acctatcccg 300
 tggccatccg ctgtgactgc ggagcctgct ccactgccac cacggagtgt gagaccatct 360
 gaggatccgg gcccaagggc gaattcgcgg ccgctaaatt caattcgccc tatagtgagt 420
 <210> 85
 <211> 131
 <212> PRT
 <213> Artificial Sequence
<220>
 <223> Description of Artificial Sequence: Fusion Protein
<400> 85
Leu Glu Pro Tyr Thr Ala Cys Asp Leu Ala Pro Pro Ala Gly Thr Thr
Asp Ala Ala His Pro Gly Tyr Leu Glu Ala Ser Ser Gly Asn Leu Arg
Thr Phe Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys
Pro Gly Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg
Cys Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala
His His Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys
Leu Pro Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val
Ala Ile Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys
        115
                            120
                                                 125
Glu Thr Ile
    130
<210> 86
<211> 420
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Fusion Protein
<400> 86
cetggagece tacacegeet gegacetgge geceeeegee ggcaceaeeg acgeegegea 60
cccgggttat ctcgaggcct ccagtgggaa cctgcgcacc tttgtgggct gtgccgtgag 120
ggagtttact ttcctggcca agaagccagg ctgcaggggc cttcggatca ccacggatgc 180
ctgctggggt cgctgtgaga cctgggagaa acccattctg gaacccccct atattgaagc 240
ccatcatcga gtctgtacct acaacgagac caaacaggtg actgtcaagc tgcccaactg 300
tgccccggga gtcgacccct tctacaccta tcccgtggcc atccgctgtg actgcggagc 360
ctgctccact gccaccacgg agtgtgagac catctgagga tccgggcccg aacaaaaact 420
```

<210> 87

<211> 387

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Fusion Protein

<400> 87

Met Lys Leu Ala Phe Leu Phe Leu Gly Pro Met Ala Leu Leu Leu 1 5 10 15

Ala Gly Tyr Gly Cys Val Leu Gly Ala Ser Ser Gly Asn Leu Arg Thr
20 25 30

Phe Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys Pro 35 40 45

Gly Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg Cys
50 60

Glu Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His 65 70 75 80

His Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys Leu

85 90 • 95

Pro Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val Ala 100 105 110

Ile Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys Glu
115 120 125

Thr:Ile Asp Lys Gly Gln Phe Cys Arg Tyr Pro Ala Gln Trp Arg Pro 130 135 140

Leu Glu Ser Arg Met Ala Ser Lys Gly Glu Glu Leu Phe Thr Gly Val 145 150 155 160

Val Pro Ile Leu Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe 165 170 175

Ser Val Ser Gly Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr 180 185 190

Leu Lys Phe Ile Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr 195 200 205

Leu Val Thr Thr Phe Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro 210 215 220

Asp His Met Lys Arg His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly 225 230 235 240

Tyr Val Gln Glu Arg Thr Ile Ser Phe Lys Asp Asp Gly Asn Tyr Lys 245 250 255

Thr Arg Ala Glu Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile 260 265 270

```
Glu Leu Lys Gly Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His
         275
Lys Leu Glu Tyr Asn Tyr Asn Ser His Asn Val Tyr Ile Thr Ala Asp
                         295
Lys Gln Lys Asn Gly Ile Lys Ala Asn Phe Lys Ile Arg His Asn Ile
 305
                                         315
Glu Asp Gly Ser Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro
                 325
Ile Gly Asp Gly Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr
                                 345
Gln Ser Ala Leu Ser Lys Asp Pro Asn Glu Lys Arg Asp His Met Val
                             360
Leu Leu Glu Phe Val Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu
                        375
Leu Tyr Lys
385
:<210> 88
<211> 1210
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Fusion Protein
<400> 88
gcatgaagct ggcattcctc ttccttggcc ccatggccct cctccttctg gctggctatg 60
gctgtgtcct cggtgcctcc agtgggaacc tgcgcacctt tgtgggctgt gccgtgaggg 120
agtttacttt cctggccaag aagccaggct gcaggggcct tcggatcacc acggatgcct 180
gctggggtcg ctgtgagacc tgggagaaac ccattctgga accccctat attgaagccc 240
atcatcgagt ctgtacctac aacgagacca aacaggtgac tgtcaagctg cccaactgtg 300
ccccgggagt cgaccccttc tacacctatc ccgtggccat ccgctgtgac tgcggagcct 360
gctccactgc caccacggag tgtgagacca tcgataaagg gcaattctgc agatatccag 420
cacagtggcg gccgctcgag tctagaatgg ctagcaaagg agaagaactt ttcactggag 480
ttgtcccaat tcttgttgaa ttagatggtg atgttaatgg gcacaaattt tctgtcagtg 540
gagagggtga aggtgatgct acatacggaa agcttaccct taaatttatt tgcactactg 600
gaaaactacc tgttccatgg ccaacacttg tcactacttt ctcttatggt gttcaatgct 660
tttcccgtta tccggatcat atgaaacggc atgacttttt caagagtgcc atgcccgaag 720
gttatgtaca ggaacgcact atatctttca aagatgacgg gaactacaag acgcgtgctg 780
aagtcaagtt tgaaggtgat accettgtta atcgtatcga gttaaaaggt attgatttta 840
aagaagatgg aaacattete ggacacaaac tegagtacaa etataaetea cacaatgtat 900
acatcacggc agacaaacaa aagaatggaa tcaaagctaa cttcaaaatt cgccacaaca 960
ttgaagatgg atccgttcaa ctagcagacc attatcaaca aaatactcca attggcgatg 1020
gccctgtcct tttaccagac aaccattacc tgtcgacaca atctgccctt tcgaaagatc 1080
ccaacgaaaa gcgtgaccac atggtccttc ttgagtttgt aactgctgct gggattacac 1140
atggcatgga tgagctctac aaataatgaa ttaaacccgc tgatcagcct cgactgtgcc 1200
ttctagttgc
<210> 89
<211> 129
```

<212> PRT

```
<213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Fusion Protein
 <400> 89
Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala Asp
                                      10
Tyr Lys Asp Asp Asp Lys Ala Ser Ser Gly Asn Leu Arg Thr Phe
Val Gly Cys Ala Val Arg Glu Phe Thr Phe Leu Ala Lys Lys Pro Gly
Cys Arg Gly Leu Arg Ile Thr Thr Asp Ala Cys Trp Gly Arg Cys Glu
                          55
Thr Trp Glu Lys Pro Ile Leu Glu Pro Pro Tyr Ile Glu Ala His His
Arg Val Cys Thr Tyr Asn Glu Thr Lys Gln Val Thr Val Lys Leu Pro
Asn Cys Ala Pro Gly Val Asp Pro Phe Tyr Thr Tyr Pro Val Ala Ile
                                 105
Arg Cys Asp Cys Gly Ala Cys Ser Thr Ala Thr Thr Glu Cys Glu Thr
                            120
Ile
<210> 90
<211> 490.
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Fusion Protein
<400> 90
ccgttgacgc aaatgggcgg taggcgtgta cggtgggagg tctatataag cagagctcgt 60
ttagtgaacc gtcagaatta attcaccatg tctgcacttc tgatcctagc tcttgttgga 120
gctgcagttg ctgactacaa agacgatgac gacaaggcct ccagtgggaa cctgcgcacc 180
tttgtgggct gtgccgtgag ggagtttact ttcctggcca agaagccagg ctgcaggggc 240
cttcggatca ccacggatgc ctgctggggt cgctgtgaga cctgggagaa acccattctg 300
gaacccccct atattgaagc ccatcatcga gtctgtacct acaacgagac caaacaggtg 360
actgtcaagc tgcccaactg tgccccggga gtcgacccct tctacaccta tcccgtggcc 420
atccgctgtg actgcggagc ctgctccact gccaccacgg agtgtgagac catctgagga 480
tcccgggtgg
<210> 91
<211> 129
<212> PRT
<213> Homo sapiens
<400> 91
Met Pro Met Ala Ser Pro Gln Thr Leu Val Leu Tyr Leu Leu Val Leu
```

1 5 10 15 Ala Val Thr Glu Ala Trp Gly Gln Glu Ala Val Ile Pro Gly Cys His Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu 105 Ile Leu Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg 120 Tyr <210> 92 <211> 490 <212> DNA <213> Homo sapiens <400> 92 ggcgaattgg gtaccgggcc cccctcgag gtcgacggta tcgataagct tagtgatgcc 60 tatggcgtcc cctcaaaccc tggtcctcta tctgctggtc ctggcagtca ctgaagcctg 120 gggccaggag gcagtcatcc caggctgcca cttgcacccc ttcaatgtga cagtgcgaag 180 tgaccgccaa ggcacctgcc agggctccca cgtggcacag gcctgtgtgg gccactgtga 240 gtccagcgcc ttcccttctc ggtactctgt gctggtggcc agtggttacc gacacaacat 300 cacctccgtc tetcagtgct gcaccatcag tggcctgaag aaggtcaaag tacagctgca 360 gtgtgtgggg agccggaggg aggagctcga gatcttaacg gccagggcct gccagtgtga 420 catgtgtcgc ctctctcgct acgaattcct gcagcccggg ggatccacta gttctagagc 480 ggccgccacc 490 · <210> 93 <211> 129 <212> PRT <213> Homo sapiens <400> 93 Met Pro Met Ala Ser Pro Gln Thr Leu Val Leu Tyr Leu Leu Val Leu Ala Val Thr Glu Ala Trp Gly Gln Glu Ala Val Ile Pro Gly Cys His Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys 35 Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser 55

Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His 70 Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu 105 Ile Leu Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg 120 Tyr <210> 94 <211> 390 <212> DNA <213> Homo sapiens <400> 94 atgcctatgg cgtcccctca aaccctggtc ctctatctgc tggtcctggc agtcactgaa 60 gcctggggcc aggaggcagt catcccaggc tgccacttgc accccttcaa tgtgacagtg 120 cgaagtgacc gccaaggcac ctgccagggc tcccacgtgg cacaggcctg tgtgggccac 180 tgtgagtcca gcgccttccc ttctcggtac tctgtgctgg tggccagtgg ttaccgacac 240 aacatcacct ccgtctctca gtgctgcacc atcagtggcc tgaagaaggt caaagtacag 300 ctgcagtgtg tggggagccg gagggaggag ctcgagatct taacggccag ggcctgccag 360 tgtgacatgt gtcgcctctc tcgctactag 390 <210> 95 <211>. 129 <212> PRT <213> Homo sapiens <400> :95 Met Pro Met Ala Ser Pro Gln Thr Leu Val Leu Tyr Leu Leu Val Leu 10 Ala Val Thr Glu Ala Trp Gly Gln Glu Ala Val Ile Pro Gly Cys His 25 Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys 40 Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser 55 Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His 65 70 Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys 90 Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu 100

125

Ile Phe Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg

120

115

490

Tyr

```
<210> 96<sup>-</sup>
 <211> 490
 <212> DNA
 <213> Homo sapiens
 <400> 96
 ggcgaattgg gtaccgggcc cccctcgag gtcgacggta tcgataagct tagtgatgcc 60
 tatggcgtcc cctcaaaccc tggtcctcta tctgctggtc ctggcagtca ctgaagcctg 120
 gggccaggag gcagtcatcc caggctgcca cttgcacccc ttcaatgtga cagtgcgaag 180
 tgaccgccaa ggcacctgcc agggctccca cgtggcacag gcctgtgtgg gccactgtga 240
 gtccagcgcc ttcccttctc ggtactctgt gctggtggcc agtggttacc gacacaacat 300
 cacctccgtc tctcagtgct gcaccatcag tggcctgaag aaggtcaaag tacagctgca 360
 gtgtgtgggg agccggaggg aggagctcga gatcttcacg gccagggcct gccagtgtga 420
 catgtgtcgc ctctctcgct acgaattcct gcagcccggg ggatccacta gttctagagc 480
 ggccgccacc
 <210> 97
 <211> 386
<212> PRT
<213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Fusion Protein
 <400> 97
 Met Pro Met Ala Ser Pro Gln Thr Leu Val Leu Tyr Leu Leu Val Leu
                                       10
 Ala Val Thr Glu Ala Trp Gly Gln Glu Ala Val Ile Pro Gly Cys His
Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys
                              40
Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser
                          55
Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His
                      70
Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys
                  85
                                      90
Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu
                                 105
Ile Leu Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg
        115
                             120
Tyr Glu Phe Cys Ser Arg Arg Tyr Arg Gly Pro Gly Ile His Arg Pro
                         135
Val Ala Thr Met Val Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val
145
                     150
                                         155
                                                              160
```

----

```
Pro Ile Leu Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser
                                     170
 Val Ser Gly Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu
                                 185
 Lys Phe Ile Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu
                             200
 Val Thr Thr Leu Thr Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp
                         215
His Met Lys Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr
225
Val Gln Glu Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr
                                     250
Arg Ala Glu Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu
                                 265
Leu Lys Gly Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys
Leu Glu Tyr Asn Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys
                         295
Gln Lys Asn Gly Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu
305
                     310
                                         315
Asp Gly Ser Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile
                                     330
Gly Asp Gly Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln
                                 345
Ser Ala Leu Ser Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu
        355
                             360
Leu Glu Phe Val Thr Ala Ala Gly Ile Thr Leu Gly Met Asp Glu Leu
                        375
Tyr Lys
385
<210> 98
<211> 1190
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Fusion Protein
<400> 98
agettagtga tgcctatggc gtcccctcaa accetggtcc tctatctgct ggtcctggca 60
gtcactgaag cctggggcca ggaggcagtc atcccaggct gccacttgca ccccttcaat 120
gtgacagtgc gaagtgaccg ccaaggcacc tgccagggct cccacgtggc acaggcctgt 180
gtgggccact gtgagtccag cgccttccct tctcggtact ctgtgctggt ggccagtggt 240
taccgacaca acatcacctc cgtctctcag tgctgcacca tcagtggcct gaagaaggtc 300
aaagtacage tgcagtgtgt ggggageegg agggaggage tegagatett aaeggeeagg 360
```

```
gcctgccagt gtgacatgtg tcgcctctct cgctacgaat tctgcagtcg acggtaccgc 420
  gggcccggga tccaccggcc ggtcgccacc atggtgagca agggcgagga gctgttcacc 480
  ggggtggtgc ccatcctggt cgagctggac ggcgacgtaa acggccacaa gttcagcgtg 540
  teeggegagg gegagggega tgecacetae ggeaagetga eeetgaagtt catetgeace 600
  aceggeaage tgeeegtgee etggeeeace etegtgaeea eeetgaeeta eggegtgeag 660
  tgcttcagcc gctaccccga ccacatgaag cagcacgact tcttcaagtc cgccatgccc 720
  gaaggctacg tccaggagcg caccatcttc ttcaaggacg acggcaacta caagacccgc 780
  gccgaggtga agttcgaggg cgacaccctg gtgaaccgca tcgagctgaa gggcatcgac 840
  ttcaaggagg acggcaacat cctggggcac aagctggagt acaactacaa caqccacaac 900
  gtctatatca tggccgacaa gcagaagaac ggcatcaagg tgaacttcaa gatccgccac 960
  aacatcgagg acggcagcgt gcagctcgcc gaccactacc agcagaacac ccccatcggc 1020
  gacggccccg tgctgctgcc cgacaaccac tacctgagca cccagtccgc cctgagcaaa 1080
  gaccccaacg agaagcgcga tcacatggtc ctgctggagt tcgtgaccgc cgccgggatc 1140
  actctcggca tggacgagct gtacaagtaa agcggccgcg actctagatc
  <210> 99
  <211> 165
 <212> PRT
  <213> Artificial Sequence
<220>
  <223> Description of Artificial Sequence: Fusion Protein
 <400> 99
 Ala Ala Cys Leu Glu Pro Tyr Thr Ala Cys Asp Leu Ala Pro Pro Ala
   1
                                       10
 Gly Thr Thr Asp Ala Ala His Pro Gly Tyr Leu Glu Glu Ala Leu Ser
 Leu Glu Gln Glu Ala Val Ile Pro Gly Cys His Leu His Pro Phe Asn
 Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys Gln Gly Ser His Val
 Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser Ala Phe Pro Ser Arg
 Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His Asn Ile Thr Ser Val
 Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys Val Lys Val Gln Leu
 Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu Ile Phe Thr Ala Arg
 Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg Tyr Glu Phe Gly Pro
 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser Ala Val Asp His
 145
                                         155
                                                              160
 His His His His
```

<210> 100 <211> 560 165

```
<212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Fusion Protein
 <400> 100
 geegeetgee tggageeeta caeegeetge gaeetggege ceeeegeegg caeeaeegae 60
 gccgcgcacc cgggttatct cgaggaagcg ctctctctag aacaggaggc agtcatccca 120
 ggctgccact tgcacccctt caatgtgaca gtgcgaagtg accgccaagg cacctgccag 180
 ggctcccacg tggcacaggc ctgtgtgggc cactgtgagt ccagcgcctt cccttctcgg 240
 tactctgtgc tggtggccag tggttaccga cacaacatca cctccgtctc tcagtgctgc 300
 accatcagtg gcctgaagaa ggtcaaagta cagctgcagt gtgtggggag ccggagggag 360
 gagetegaga tetteaegge eagggeetge eagtgtgaea tgtgtegeet etetegetae 420
gaattcgggc ccgaacaaaa actcatctca gaagaggatc tgaatagcgc cgtcgaccat 480
 catcatcatc atcattgagt ttaaacccgc tgatcagcct cgactgtgcc ttctagttgc 540
cagccatctg ttgtttgccc
 <210> 101
 <211> 129
 <212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Fusion Protein
<400> 101
Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala Asp
Tyr Lys Asp Asp Asp Lys Gln Glu Ala Val Ile Pro Gly Cys His
                                  25
Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg Gln Gly Thr Cys
Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His Cys Glu Ser Ser
                         55
Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser Gly Tyr Arg His
Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser Gly Leu Lys Lys
Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg Glu Glu Leu Glu
Ile Phe Thr Ala Arg Ala Cys Gln Cys Asp Met Cys Arg Leu Ser Arg
                            120
                                                125
Tyr
<210> 102
<211> 420
<212> DNA
<213> Artificial Sequence
```

```
<220>
 <223> Description of Artificial Sequence: Fusion Protein
 agetgetage caccatgtet geacttetga tectagetet tgttggaget geagttgetg 60
 actacaaaga cgatgacgac aagcaggagg cagtcatccc aggctgccac ttgcacccct 120
 tcaatgtgac agtgcgaagt gaccgccaag gcacctgcca gggctcccac gtggcacagg 180
 cetgtgtggg ccactgtgag tecagegeet tecetteteg gtactetgtg etggtggeea 240
gtggttaccg acacaacatc acctccgtct ctcagtgctg caccatcagt ggcctgaaga 300
aggtcaaagt acagctgcag tgtgtgggga gccggaggga ggagctcgag atcttcacgg 360
ccagggcctg ccagtgtgac atgtgtcgcc tctctcgcta ctgaggatcc agacatgata 420
 <210> 103
 <211> 69
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 103
ctcttgttgg agctgcagtt gctcatcatc accatcacca tggtgacgat gacgataagc 60
aggaggcag
<210> 104
<211> 39
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
<400> 104
tttggatccg tcgactagta gcgagagagg cgacacatg
                                                                    39
<210> 105
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: PCR Primer
      Sequence
tttgctagcg tcgaccatgt ctgcacttct gatcctagct cttgttggag ctgcagttgc 60
tcatc
                                                                   65
<210 > 106
<211> 39
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: PCR Primer
```

Sequence

<400> 106 tttggatccg tcgactagta gcgagagag cgacacatg

39

<210> 107

<211> 133

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Fusion Protein

<400> 107

Met Ser Ala Leu Leu Ile Leu Ala Leu Val Gly Ala Ala Val Ala His

1 10 15

His His His His Gly Asp Asp Asp Lys Gln Glu Ala Val Ile
20 25 30

Pro Gly Cys His Leu His Pro Phe Asn Val Thr Val Arg Ser Asp Arg 35 40 45

Gln Gly Thr Cys Gln Gly Ser His Val Ala Gln Ala Cys Val Gly His
50 60

Cys Glu Ser Ser Ala Phe Pro Ser Arg Tyr Ser Val Leu Val Ala Ser 65 70 75 80

Gly Tyr Arg His Asn Ile Thr Ser Val Ser Gln Cys Cys Thr Ile Ser 85 90 95

Gly Leu Lys Lys Val Lys Val Gln Leu Gln Cys Val Gly Ser Arg Arg 100 105 110

Glu Glu Leu Glu Ile Phe Thr Ala Arg Ala Cys Gln Cys Asp Met Cys 115 120 125

Arg Leu Ser Arg Tyr 130